

IDS  
AL

1: AA234924. zr78g10.r1 Soares...[gi:1859416] [Links](#)

dbEST Id: 874704  
EST name: zr78g10.r1  
GenBank Acc: AA234924  
GenBank gi: 1859416  
GDB Id: 5563537

Clone Id: IMAGE:669570 (5')  
Source: IMAGE Consortium, LLNL  
Insert length: 655  
DNA type: cDNA

Sequencing: -28ml3 rev2 ET from Amersham  
PolyA Tail: Unknown

GTTTTTGAGGAGCGACTGAGGCAAATGGGGTCTGGCTAAAAGTCAATGGAGAAGCTATT  
 TATGAAACCCATACCTGGCGATCCCAGAATGACACTGTCCCCAGATGTGTGGTACACAT  
 CCAAGCCTAAAGAAAAATTAGTCTATGCCATTTTTCTTAAATGGCCACATCAGGACAGC  
 TGTTCCCTTGGCCATCCCAAAGCTATTCTGGGGCAACAGAGGTGAAACTACTGGCCATG  
 GACAGCCACTTAACTGGATTCTTTGGAGCAAAATGGCATTATGGTAGAACTGCCACAGC  
 TAACCATTCATCAGATGCCGTGTAAATGGGGCTGGGCTCTAGCCTGACTAATGTGATCTA  
 AAGTGCAGCAGATGGCTGATGCTGCAAGTTATGTCTAAGGCTAGGAACATCAGGTGTC  
 TATAATTGTAGCACATGGAGAAAGCAAATGTAAACTGGATAAGAAAAAT  
 Quality: High quality sequence stops at base: 430

Entry Created: Mar 3 1997  
Last Updated: Aug 6 1997

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

PUTATIVE ID Assigned by submitter  
SW:FUCO\_RAT P17164 ALPHA-L-FUCOSIDASE PRECURSOR ;

Lib Name: Soares\_NhHMPu\_S1  
Organism: Homo sapiens  
Organ: mixed (see below)  
Tissue type: Pooled human melanocyte, fetal heart, and pregnant uterus  
Lab host: DH10B  
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker  
R. Site 1: Not I  
R. Site 2: Eco RI  
Description: Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart

NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

**SUBMITTER**

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E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

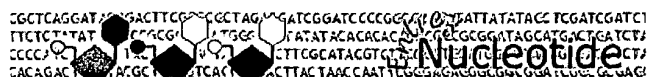
Title: WashU-Merck EST Project 1997  
Authors: Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R., Wilson, R.  
Year: 1997  
Status: Unpublished

**MAP DATA**

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Revised: July 5, 2002.

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[NCBI](#) | [NLN](#) | [NIH](#)



A H

1: W52490. zc54a02.r1 Soares...[gi:1350088]

## IDENTIFIERS

### CLONE INFO

## PRIMERS

## SEQUENCE

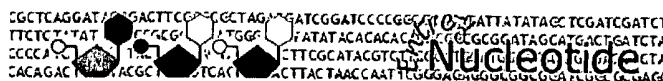
Quality: High quality sequence stops at base: 440

### COMMENTS

PUTATIVE ID

LIBRARY

1 of 2



IDS  
AJ

1: AA065306. a03500s Testis 5 ...[gi:1929306] Links

dbEST Id: 691225  
EST name: a03500s  
GenBank Acc: AA065306  
GenBank qi: 1929306

Clone Id: a03500 (3' end)  
DNA type: cDNA

Sequencing: SP6  
PolyA Tail: Unknown

AGCTAAGAAAGAACTTTATTATTTCTAATAAATGCATCTGGGGGTGGGGTCTGGAGAGGG  
CACACGGCCGTCCCCTGCCAATGCGTTGAGGCCCACACTCCTTGGGAGTGGACAGGGAAA  
GCCCCATGTTCCCGGAGCCTGTCGCCGTGGCCTGCAGCCCTGGACGCCTNTNACGGAAC  
CGCAGTCTTTCTCCCGGCTTTCTGGTGCCTGGCCACCACAGCCCCATCAATACAGCC  
AGAGCCCGGGGCCAGGTGTGAGTNTGCCGTAGCTNTCCGTCTCCGNCAGCCCCGACGNN  
TTGGAGCAGNTTCTGAAGGCCTTCTGGGTGTTAGCATCCGTTGGGGAGCGGGTTTTTGC  
GTGTTGGGA

Entry Created: Dec 31 1996  
Last Updated: Jun 12 2002

This sequence derives from a clone which was selected from the cDNA library - Testis 5 - using a repeat of 14 CAG as probe

```

Lib Name:      Testis 5
Organism:      Homo sapiens
Vector:        pSPORT1
R. Site 1:     MluI
R. Site 2:     NotI
Description:    mRNA was prepared from human testis of a 27 years old man.
                cDNA was prepared using a 15mer oligo dT anchored by two
                degenerated bases at its 3'end and containing a NotI site at
                its 5'end. The cDNA was cloned between SalI and NotI sites
                of pSPORT1. The MluI-SalI fragement come from the adaptator
                used for the cloning. The 3' end is at the NotI site. cDNA
                corresponding to abundant species were eliminated from this
                library.

```

Name: Guellaen G  
Lab: Unite INSERM 99  
Institution: INSERM

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Fax: (33)148980908  
E-mail: [guellaen@infobiogen.fr](mailto:guellaen@infobiogen.fr)

**CITATIONS**

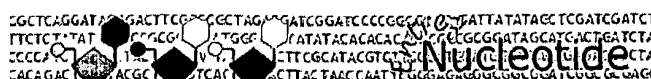
Title: Guellaen, G. Unpublished (1996)  
Authors: Guellaen, G.  
Year: 1996  
Status: Unpublished

**MAP DATA**

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Revised: July 5, 2002.

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IDS  
AI

□1: AA065307. a03500t Testis 5 ...[gi:1929307]

## Links

dbEST Id: 691226  
EST name: a03500t  
GenBank Acc: AA065307  
GenBank gi: 1929307

Clone Id: a03500 (5' end)  
DNA type: cDNA

Sequencing: T7  
PolyA Tail: Unknown

GGAAATCTAACAAATCCCTGGGGAGACCGGAGACCGGTGACCGCCGCCCTGCCCGCAGCCATG  
TGGCCCCCGCTGTGCTGCTGCTGCTGCTCCCGGCCGCCCGGTCCCACCGCCAAA  
GCCGCTCCCCACCCGGATGCTAACACCCAGGAAGGCCCTNAGAACCCTGCTCCAAGGCGTC  
GGGGCTGGNGGAGACGNAGAGTGCGGGCAGACTNACACTTGCCCCCGGGCTCTGGCTGT  
ATTGATGGGGCTGTGGTGGCCACGTGACACAGAAAGCNGNGGAGGAAGACC

Entry Created: Dec 31 1996  
Last Updated: Jun 12 2002

This sequence derives from a clone which was selected from the cDNA library - Testis 5 - using a repeat of 14 CAG as probe

```

Lib Name:      Testis 5
Organism:      Homo sapiens
Vector:        pSPORT1
R. Site 1:     MluI
R. Site 2:     NotI
Description:    mRNA was prepared from human testis of a 27 years old man.
                cDNA was prepared using a 15mer oligo dT anchored by two
                degenerated bases at its 3'end and containing a NotI site at
                its 5'end. The cDNA was cloned between Sall and NotI sites
                of pSPORT1. The MluI-Sall fragement come from the adaptator
                used for the cloning. The 3' end is at the NotI site. cDNA
                corresponding to abundant species were eliminated from this
                library.

```

Name: Guellaen G  
Lab: Unite INSERM 99  
Institution: INSERM  
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Tel: (33)149813530

Fax: (33)148980908  
E-mail: [guellaen@infobiogen.fr](mailto:guellaen@infobiogen.fr)

**CITATIONS**

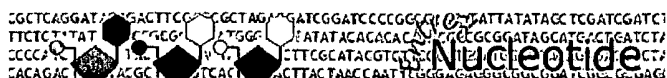
Title: Guellaen, G. Unpublished (1996)  
Authors: Guellaen, G.  
Year: 1996  
Status: Unpublished

**MAP DATA**

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IDS  
AK

1: AA151194.zl40b11.r1 Soares...[gi:1719402] [Links](#)

dbEST Id: 787792  
EST name: z140b11.r1  
GenBank Acc: AA151194  
GenBank gi: 1719402  
GDB Id: 3809369

Clone Id: IMAGE:504381 (5')  
Source: IMAGE Consortium, LLNL  
Insert length: 889  
DNA type: cDNA

Sequencing: -28M13 rev2 from Amersham  
PolyA Tail: Unknown

ATTCGGCACGACGGTGATCGTTATAACCCAGGACATCTTTTGCCACATAAATGGGAAAAAC  
 TGCATGACAATAGACAACTGTCTGGGGCTATAGGAGGGAAGCTGGAATCTCTGACTAT  
 CTTACAATTGAAGAATTGGTGAAGCAACTTGTAGAGACAGTTTCATGTGGAGGAAATCTT  
 TTGATGAATATTGGGCCACACTAGATGGCACCATTTCGTAGTTTTGGAGGAGCGACTG  
 AGGCAAGTGGGGTCTCTGGCTAAAAGTCAATGGAGAAGCTATTTATGAAACCTATACCTGG  
 CGATCCCAGAATGACACTGTCACCCCAGATGTGTGGTACACATCCAAGCCTAAAGAAAAA  
 TTAGTCTATGCCATTTTCTTAAATGGCCCACATCAGGACAGCTGTTCTTGGCCATCCC  
 AAAGCTATCTGGGGGCAACAGAGGTGAAACCTACTGGGCCATGGAG  
 Quality: High quality sequence stops at base: 435

Quality: High quality sequence stops at base: 435

Entry Created: Nov 15 1996  
Last Updated: May 14 1997

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

PUTATIVE ID Assigned by submitter  
TR:G182779 G182779 LYSOSOMAL ENZYME ALPHA-L-FUCOSIDASE ;

```

Lib Name:      Soares_pregnant_uterus_NbHPU
Organism:      Homo sapiens
Sex:           female
Organ:          uterus
Develop. stage: adult
Lab host:       DH10B
Vector:         pT7T3-Pac
R. Site 1:      Not I
R. Site 2:       Eco RI
Description:    1st strand cDNA was primed with a Not I - oligo(dT) primer

```